



PATENT
Attorney Docket No. 8959.0009

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:)	
Katsura Nozawa et al.)	Group Art Unit: 1649
Application No.: 10/506,624)	Examiner: Kimberly A. Ballard, Ph.D.
Filed: September 3, 2004)	
For: NOVEL POTASSIUM-DEPENDENT SODIUM-CALCIUM EXCHANGER)	Confirmation No.: 1864

Commissioner for Patents
P.O. Box 1450
Alexandria, VA 22313-1450

Sir:

**SUBMISSION OF CERTIFIED ENGLISH TRANSLATIONS
OF PRIORITY DOCUMENTS**

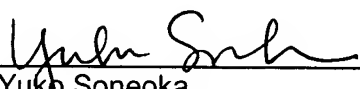
Enclosed herewith are certified English translations of Japanese priority documents JP 2002-225114, filed August 1, 2002, and JP 2003-182989, filed June 26, 2003. The attached Certificates of Translation include a statement from the translator that the English translations are accurate. In accordance with 37 C.F.R. § 1.55(a), Applicants hereby perfect claim of priority under 35 U.S.C. § 119 by filing these certified translations.

Please grant any extensions of time required to enter this response and charge any additional required fees to our deposit account 06-0916.

Respectfully submitted,

FINNEGAN, HENDERSON, FARABOW,
GARRETT & DUNNER, L.L.P.

Dated: December 29, 2006

By: 
Yuko Soneoka
Limited Reg. No. L0093

CERTIFICATE OF VERIFICATION

I, Masayuki HARABAYASHI,
of c/o Morita & Associates, 5th Fl., Itabashi-Chuo Bldg.,
67-8, Itabashi 2-chome, Itabashi-ku, Tokyo 173-0004, Japan,
state that the attached document is a true and complete
translation to the best of my knowledge of Japanese Patent
Application No. 2003-182989.

Dated this 1st day of December, 2006

Signature of translator: M. Harabayashi

[DOCUMENT NAME] Patent Application
[REFERENCE NUMBER] YAM033286P
[ADDRESSEE] Director-General of Patent Office
[INTERNATIONAL PATENT CLASSIFICATION] C07K 14/00
[INVENTOR]
 [Address or Residence]
 c/o Yamanouchi Pharmaceutical Co., Ltd.,
 21, Miyukigaoka, Tsukuba-shi, Ibaraki 305-8585, Japan
 [Name] Katsura NOZAWA
[INVENTOR]
 [Address or Residence]
 c/o Yamanouchi Pharmaceutical Co., Ltd.,
 21, Miyukigaoka, Tsukuba-shi, Ibaraki 305-8585, Japan
 [Name] Shinobu MOCHIZUKI
[APPLICANT]
 [Identification No.] 000006677
 [Name] Yamanouchi Pharmaceutical Co., Ltd.
[ATTORNEY]
 [Identification No.] 100090251
 [Patent Attorney]
 [Name] Kenichi MORITA
[SELECTED ATTORNEY]
 [Identification No.] 100089200
 [Patent Attorney]
 [Name] Shozo NAGAI
[SELECTED ATTORNEY]
 [Identification No.] 100098501
 [Patent Attorney]
 [Name] Taku MORITA
[SELECTED ATTORNEY]
 [Identification No.] 100109357
 [Patent Attorney]
 [Name] Emiko YANO
[INDICATION OF FEE]
 [Deposit Account Number] 017813
 [Fee] 21000
[LIST OF SUBMITTED DOCUMENTS]
 [Name of Document] Specification 1
 [Name of Document] Abstract 1
[REQUEST FOR A PROOF] Yes

[DOCUMENT NAME] SPECIFICATION

[TITLE OF THE INVENTIONS] METHOD OF SCREENING THERAPEUTIC AGENTS FOR ISCHEMIA-REPERFUSION INJURY AND INFLAMMATORY DISEASES

[CLAIMS]

[Claim 1] (1) A polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 4, or (2) a polypeptide exhibiting a potassium-dependent sodium-calcium exchange activity and consisting of an amino acid sequence in which 1 to 5 amino acids in total are substituted, deleted, inserted, and/or added at one or plural portions in the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 4.

[Claim 2] A polynucleotide encoding the polypeptide according to claim 1.

[Claim 3] An expression vector comprising the polynucleotide according to claim 2.

[Claim 4] A cell transfected with the expression vector according to claim 3.

[Claim 5] A method for producing the polypeptide according to claim 1, characterized by using the cell according to claim 4.

[Claim 6] A method for screening an inhibitor of the polypeptide according to claim 1, comprising the steps of:
(1) bringing a cell expressing the polypeptide into contact with a substance to be tested,
(2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and
(3) selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide.

[Claim 7] A method for screening an inhibitor of leukocyte activation, comprising the steps of:

(1) bringing a cell expressing the polypeptide according to claim 1 into contact with a substance to be tested,
(2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and
(3) selecting a substance which inhibits the potassium-

dependent sodium-calcium exchange activity in the polypeptide.

[Claim 8] A method for screening a therapeutic agent for postischemic reperfusion injury and/or an inflammatory disease, comprising the steps of:

- (1) bringing a cell expressing the polypeptide according to claim 1 into contact with a substance to be tested,
- (2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and
- (3) selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide.

[Claim 9] A process for manufacturing a pharmaceutical composition for treating postischemic reperfusion injury and/or an inflammatory disease, comprising the steps of:

- (1) bringing a cell expressing the polypeptide according to claim 1 into contact with a substance to be tested,
- (2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and
- (3) preparing a medicament containing the substance.

[Claim 10] A pharmaceutical composition for inhibiting leukocyte activation, comprising as an active ingredient a substance obtainable by the method according to claim 6.

[Claim 11] A pharmaceutical composition for treating postischemic reperfusion injury and/or an inflammatory disease, comprising as an active ingredient a substance obtainable by the method according to claim 6.

[DETAILED DESCRIPTION OF THE INVENTION]

[0001]

[Technical Field to which the Invention Pertains]

This invention relates to a potassium-dependent sodium-calcium exchanger useful in screening an agent for treating a cell injury due to postischemic reperfusion and/or an inflammatory disease.

[0002]

[Prior Art]

During ischemia, an organ proceeds via a reversible to

the irreversible phase of cell injury. It is considered that an injury can be avoided if the organ is reperfused in the process, but experimental and clinical examinations point out that the reperfusion per se newly damages the organ. This is called a reperfusion injury. It is known that both free radicals generated during reperfusion and excess intracellular calcium (calcium overload) due to reperfusion play an important role in the generation of reperfusion injuries (non-patent references 1 and 2). Regarding the mechanism of the calcium overload, several hypotheses are proposed, as follows (non-patent reference 3):

(1) Due to a failure of an energy-dependent calcium exchange system, calcium flows into cells in accordance with a concentration gradient thereof.

(2) A concentration of intracellular sodium ions is increased during ischemia, and therefore calcium flows into cells via a sodium-calcium exchange system.

(3) An increased α receptor density during ischemia promotes a calcium overload after reperfusion.

(4) Due to a severe failure of a cell membrane, extracellular calcium flows into cells in accordance with a concentration gradient thereof.

Further, it is considered that, in delayed neuronal cell death during postischemic reperfusion in the brain, a glutamate receptor, NMDA (N-methyl-D-aspartate) receptor, is activated, an NMDA activated calcium channel is opened, and thus calcium flows into cells (non-patent reference 4).

[0003]

After transient ischemia in an experimental animal, the phenomenon of no reflow (when a blood flow is reflowed, a region where a blood flow is not topically reflowed is generated) is observed in the brain, heart, and kidney. As major mechanisms of the phenomenon of no reflow, hemorrhage and destruction of microvascular construction (for example, capillary embolus by leukocytes, or swelling of vascular endothelial cells), platelet thrombosis, or the like are suggested (non-patent reference 5). As cells which block microvessels in a focus of middle cerebral artery occlusion,

polymorphonuclear leucocytes, monocytes, or platelets are observed in addition to erythrocytes, and particularly, capillary embolus by neutrophils classified into leukocytes is noted as a cause of the phenomenon of no reflow (non-patent reference 5). When passing through microvessels, polymorphonuclear leucocytes delay an erythrocyte flow. Further, during ischemia in the brain, polymorphonuclear leucocytes interact with vascular endothelial cells or platelets, and expressions of receptors for adhesion factors are increased at the surfaces of polynuclear leucocytes. Furthermore, monoclonal antibodies against adhesion factors such as L-selection, CD11, CD18, or ICAM-1 attenuated leukocyte accumulation and reduced the area of infarction, and thus it was suggested that the expression of adhesion factors plays an important role in accumulation of leukocytes during postischemic reperfusion (non-patent references 6 and 7). In addition, it is reported that the expression of adhesion factors in leukocytes is increased by increasing an intracellular calcium concentration (non-patent reference 8).

[0004]

Numerous ion channels controlling an intracellular ion environment are known. As those which play a particularly important role in postischemic reperfusion injury, for example, a sodium-hydrogen exchanger and a sodium-calcium exchanger are known. During the process of postischemic reperfusion injury, there are two stages in the passage from ischemia to reperfusion. One is ischemia and the other is reperfusion. During ischemia, blood is not supplied and an anaerobic condition is brought about, and thus an exhaustion of energy, promotion of anaerobic metabolism, accumulation of harmful metabolites, and the like occur, and, as a result, acidosis proceeds. During reperfusion, oxygen is supplied, and thus acidosis is attenuated, together with a supply of energy, restart of aerobic metabolism, and removal of accumulated metabolites, to return the physiological conditions. However, because injury proceeds rapidly during reperfusion, it seems that the conditions neutralized by reperfusion are worse than that of acidosis by ischemia. It

is known that the sodium-hydrogen exchanger and the sodium-calcium exchanger are involved in this phenomenon (non-patent reference 9).

[0005]

It is known that the sodium-hydrogen exchanger exists on the cell membrane, is involved in a control of intracellular pH, and has an activity of extruding hydrogen ions outside the cell in accordance with a concentration gradient during acidosis, and simultaneously, incorporating sodium ions into the cell. It is reported that the mechanism in which an expression and activity of the sodium-hydrogen exchanger are suppressed during ischemic preconditioning is involved in a protective action (non-patent reference 10). Further, it is reported that the activation of the sodium-hydrogen exchanger activates neutrophils and promotes neutrophil adhesion via expression of CD11, CD18, or ICAM-1 (non-patent reference 11).

[0006]

There are two types of sodium-calcium exchange reactions, i.e., the classic sodium-calcium exchange reaction and the potassium-dependent sodium-calcium exchange reaction (non-patent references 12 and 13). It is considered that sodium-calcium exchangers exist in all tissues. The physiological function is an extrusion of intracellular calcium outside the cell, accompanied by an influx of extracellular sodium. Further, it is reported that the reverse reaction in which intracellular sodium is extruded and calcium flows into the cell may occur physiologically.

[0007]

According to a report about ion dynamics during postischemic reperfusion, oxygen is not supplied and anaerobic metabolism is promoted during ischemia, and thus the substrate (ATP) for energy is exhausted and acidosis proceeds. That is, sodium ions are not extruded outside the cell due to the inactivation of sodium pumps, and sodium ions flow into the cell by gradually extruding hydrogen ions outside the cell via the sodium-hydrogen exchanger for an adjustment of acidosis. The accumulation of intracellular

sodium ions results in an increase in intracellular calcium ions via the sodium-calcium exchanger. Further, because the pH around tissues is neutralized rapidly during reperfusion, a concentration gradient between intracellular hydrogen ions accumulated by anaerobic metabolism and decreased hydrogen ions therearound becomes large. In addition, the sodium-hydrogen exchanger is activated, and then hydrogen ions are extruded outside the cell and sodium ions rapidly flow into the cell. After the rapid influx of sodium ions, the sodium-calcium exchanger is activated, and then sodium ions are extruded outside the cell and calcium ions are transported into the cell. That is, a concentration of intracellular calcium is increased rapidly to become a calcium overload. This rapidly increased pH and calcium influx into the cell is considered to be a trigger for injurious actions occurring during postischemic reperfusion (non-patent reference 9). As described above, the activation of the sodium-hydrogen exchanger as the first step, and the activation of the sodium-calcium exchanger as the second step, play an important role in a relatively early stage of postischemic reperfusion injury.

[0008]

In connection with the increased intracellular calcium concentration in leukocytes during postischemic reperfusion, it is known that a calcium influx into the cell is important in the process of neutrophil activation caused by leukocyte mobilizing factors such as formylmethionyl-leucinyl-phenylalanine (fMLP), arachidonic acid, or leukotoriene B₄, and that the neutrophil activation is inhibited by suppressing the influx (non-patent reference 14). Further, it is reported that intracellular pH is increased by treatment with fMLP, and that inhibitors of the sodium-hydrogen exchanger suppressed increased intracellular pH and leukocyte migration (non-patent reference 15).

However, it is not known whether the inhibition of the sodium-calcium exchanger in leukocytes suppresses leukocyte activation.

[0009]

(non-patent reference 1) "Circulation", (U.S.A.), 1990, 82,

p. 723-738

- (non-patent reference 2) "Annual Review of Physiology",
(U.S.A.), 1992, 54, p. 243-256
- (non-patent reference 3) "Kokyu To Junkan", 2001, 49(1), p.
5-11
- (non-patent reference 4) "CLINICAL NEUROSCIENCE", (U.S.A.),
1999, 17(5), p. 567-569
- (non-patent reference 5) "Kokyu To Junkan", 2001, 49(1), p.
13-20
- (non-patent reference 6) "Circulation", (U.S.A.), 1993, 88,
p. 649-658
- (non-patent reference 7) "American Journal of Pathology",
(U.S.A.), 1993, 143, p. 410-418
- (non-patent reference 8) "Cell Adhesion &
Communication", (Switzerland), 1993, 1, p. 21-32
- (non-patent reference 9) Yasuo Morishita et al., "Zoki no
Kyoketsu-saikanryu-shogai - Kiso to Rinsyo (Reperfusion
Injury)", SHINDAN-TO-CHIRYOSHA, 2002, p.1-225
- (non-patent reference 10) "Circulation Research", (U.S.A.),
1999, 85, p. 723-730
- (non-patent reference 11) "Journal of Cardiovascular
Pharmacology", (U.S.A.), 2001, 37, p. 668-677
- (non-patent reference 12) "Tanpakushitsu Kakusan Koso", 1998,
43(12), p. 1555-1560
- (non-patent reference 13) "Journal of Biological Chemistry",
(U.S.A.), 1993, 268, p. 6874-6877
- (non-patent reference 14) "Biochemical & Biophysical
Research Communications]", (U.S.A.), 1981, 103, p. 227-
232
- (non-patent reference 15) "British Journal of
Pharmacology", (United Kingdom), 1998, 124, p. 627-638

[0010]

[Problems to be Solved by the Invention]

As described in PRIOR ART, with respect to the
leukocyte activation, postischemic reperfusion and
inflammation have common aspects, such as an increased
intracellular pH in leukocytes and activation of the sodium-
hydrogen exchanger thereby, and an increased intracellular
calcium concentration thereby. It is considered that the

sodium-calcium exchanger in leukocytes is involved in leukocyte activation during postischemic reperfusion injury and inflammation. That is, it is considered that an inhibition of sodium-calcium exchanger activation in leukocytes suppresses the increased intracellular calcium concentration in leukocytes and inhibits the leukocyte activation.

The object of the present invention is to provide a novel potassium-dependent sodium-calcium exchanger in leukocytes, and a novel polynucleotide encoding the polypeptide, and further to provide a convenient screening system for obtaining a substance useful as a therapeutic agent for cell injury due to postischemic reperfusion or an inflammatory disease, of which the therapeutic effect is achieved by inhibiting an activation of leukocytes.

[0011]

[Means for Solving the Problems]

The present inventors conducted intensive studies and, as a result, obtained polynucleotides encoding potassium-dependent sodium-calcium exchangers consisting of the amino acid sequences of SEQ ID NOS: 2 and 4, respectively, and abundantly expressed in leukocytes; prepared cells expressing each polypeptide; and provided screening tools for obtaining a substance useful as an inhibitor of the sodium-calcium exchangers, that is, a therapeutic agent for postischemic reperfusion injury or an inflammatory disease. Further, the present inventors constructed systems for detecting the sodium-calcium exchange activity by using the cells expressing each sodium-calcium exchanger; and provided a method for screening a substance useful as a therapeutic agent for postischemic reperfusion injury and an inflammatory disease, on the basis of inhibition of the above activity. Furthermore, the present inventors obtained inhibitors of the sodium-calcium exchangers by the above screening method; confirmed that the inhibitors inhibited migration of human peripheral polymorphonuclear leukocytes, that is, activation of leukocytes; and provided novel therapeutic agents for postischemic reperfusion injury and/or an inflammatory disease containing each inhibitor of

leukocyte activation; and thus the present invention was completed.

[0012]

The present invention relates to:

- [1] (1) a polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 4, or (2) a polypeptide exhibiting a potassium-dependent sodium-calcium exchange activity and consisting of an amino acid sequence in which 1 to 5 amino acids in total are substituted, deleted, inserted, and/or added at one or plural portions in the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 4;
- [2] a polynucleotide encoding the polypeptide of [1];
- [3] an expression vector comprising the polynucleotide of [2];
- [4] a cell transfected with the expression vector of [3];
- [5] a method for producing the polypeptide of [1], characterized by using the cell of [4];
- [6] a method for screening an inhibitor of the polypeptide of [1], comprising the steps of:
 - (1) bringing a cell expressing the polypeptide into contact with a substance to be tested,
 - (2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and
 - (3) selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide;
- [7] a method for screening an inhibitor of leukocyte activation, comprising the steps of:
 - (1) bringing a cell expressing the polypeptide of [1] into contact with a substance to be tested,
 - (2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and
 - (3) selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide;
- [8] a method for screening a therapeutic agent for postischemic reperfusion injury and/or an inflammatory

disease, comprising the steps of:

(1) bringing a cell expressing the polypeptide of [1] into contact with a substance to be tested,

(2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and

(3) selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide;

[9] a process for manufacturing a pharmaceutical composition for treating postischemic reperfusion injury and/or an inflammatory disease, comprising the steps of:

(1) bringing a cell expressing the polypeptide of [1] into contact with a substance to be tested,

(2) analyzing whether or not a potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and

(3) preparing a medicament containing the substance;

[10] a pharmaceutical composition for inhibiting leukocyte activation, comprising as an active ingredient a substance obtainable by the method of [6]; and

[11] a pharmaceutical composition for treating postischemic reperfusion injury and/or an inflammatory disease, comprising as an active ingredient a substance obtainable by the method of [6].

[0013]

The term "therapeutic agent for postischemic reperfusion injury" or "pharmaceutical composition for treating postischemic reperfusion injury" as used herein includes both an agent or pharmaceutical composition used for treating a patient suffering postischemic reperfusion injury and that preventively used for a subject having the risk of postischemic reperfusion injury. The term "therapeutic agent for an inflammatory disease" or "pharmaceutical composition for treating an inflammatory disease" as used herein includes both an agent or pharmaceutical composition used for treating a patient suffering an inflammatory disease and that preventively used for a subject having the risk of an inflammatory disease.

[0014]

[Mode for Carrying out the Invention]

The present invention will be explained in detail hereinafter.

[0015]

1. Polypeptide, polynucleotide, expression vector, and cell of the present invention

The polypeptide of the present invention includes

(1) a polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4; and

(2) a polypeptide consisting of an amino acid sequence in which 1 to 5 amino acids in total are substituted, deleted, inserted, and/or added at one or plural portions in the amino acid sequence of SEQ ID NO: 2 or 4, and exhibiting a potassium-dependent sodium-calcium exchange activity (hereinafter referred to as a variation functionally equivalent). As the polypeptide of the present invention, the polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4 is preferable.

[0016]

The term "exhibiting a sodium-calcium exchange activity" as used herein means exhibiting an exchange reaction in which intracellular calcium is extruded outside the cell and extracellular sodium flows into the cell (the forward direction), or exhibiting an exchange reaction in which intracellular sodium is extruded outside the cell and extracellular calcium flows into the cell (the reverse direction). Further, the term "exhibiting a forward sodium-calcium exchange activity" as used herein means exhibiting an exchange reaction in which intracellular calcium is extruded outside the cell and extracellular sodium flows into the cell ($\text{Na}^+_{\text{o}}\text{-Ca}^{2+}_{\text{i}}$ exchange; the forward direction). The term "exhibiting a reverse sodium-calcium exchange activity" as used herein means exhibiting an exchange reaction in which intracellular sodium is extruded outside the cell and extracellular calcium flows into the cell ($\text{Na}^+_{\text{i}}\text{-Ca}^{2+}_{\text{o}}$ exchange; the reverse direction).

[0017]

Whether or not a polypeptide (hereinafter referred to

as a test polypeptide) "exhibits a forward sodium-calcium exchange activity", that is, "exhibits an exchange reaction in which intracellular calcium is extruded outside the cell and extracellular sodium flows into the cell (the forward direction)" may be confirmed by a method known to those skilled in the art (Iwamoto T. et al., J. Biol. Chem., 271, 22391-22397, 19962). A method for confirming it is not particularly limited and, for example, the following method may be used.

Cells are transfected with an expression vector comprising a polynucleotide encoding the test polypeptide. The resulting cells are cultured in a medium containing calcium chloride (^{45}Ca chloride) to incorporate calcium ions into the cells; the cells are washed with a washing solution to remove calcium ions not incorporated; the extracellular solution is changed to an extracellular solution for measurement containing sodium ions; and a ^{45}Ca radioactivity contained in the solution is measured. When the radioactivity is detected, it may be confirmed that the test polypeptide "exhibits a forward sodium-calcium exchange activity".

[0018]

Whether or not a test polypeptide "exhibits a reverse sodium-calcium exchange activity", that is, "exhibits an exchange reaction in which intracellular sodium is extruded outside the cell and extracellular calcium flows into the cell (the reverse direction)" may be confirmed by a method known to those skilled in the art (Iwamoto T. et al., J. Biol. Chem., 271, 22391-22397, 19962). A method for confirming it is not particularly limited and, for example, the following method (preferably a method described in Example 4) may be used.

Cells are transfected with an expression vector comprising a polynucleotide encoding the test polypeptide. The resulting cells are treated with an ionophore (such as monensin) to a univalent cation, to incorporate sodium into the cells; the extracellular solution is changed to a solution containing ^{45}Ca , to exchange intracellular sodium for extracellular calcium; and an intracellular ^{45}Ca

radioactivity is measured. When the radioactivity is detected, it may be confirmed that the test polypeptide "exhibits a reverse sodium-calcium exchange activity".

[0019]

Whether or not the sodium-calcium exchange activity of the test compound is "potassium-dependent" may be confirmed by a method known to those skilled in the art (Kimura, M., J. Biol. Chem., 268, p.6874-6877, 1993; Kraev A. et al., J. Biol. Chem., 276, 23161-23172, 2001). A method for confirming it is not particularly limited but, for example, the following method (preferably a method described in Example 4) may be used. More particularly, cells are transfected with an expression vector comprising a polynucleotide encoding the test polypeptide. When measuring the sodium-calcium exchange activity of the resulting cells, a K^+ concentration in the extracellular solution is varied, and an intracellular ^{45}Ca radioactivity in each K^+ concentration is measured. When a higher sodium-calcium exchange activity is detected in the presence of K^+ , it may be confirmed that the sodium-calcium exchange activity of the test compound is "potassium-dependent".

[0020]

The polypeptides consisting of the amino acid sequences of SEQ ID NOS: 2 and 4, which are included in the polypeptide of the present invention, are novel human potassium-dependent sodium-calcium exchangers consisting of 622 and 603 amino acid residues, respectively. Each polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4 is expressed in peripheral leukocytes, as shown in Example 2.

[0021]

The variation functionally equivalent of the present invention is not particularly limited, so long as it is a polypeptide consisting of an amino acid sequence in which 1 to 5 amino acids, preferably 1 to 3 amino acids, in total are substituted, deleted, inserted, and/or added at one or plural positions (for example, 1 to 3 positions) in the amino acid sequence of SEQ ID NO: 2 or 4, and exhibiting the potassium-dependent sodium-calcium exchange activity.

Further, an origin of the variation functionally equivalent is not limited to a human.

[0022]

The variation functionally equivalent of the present invention includes, for example, not only human variations of the polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4, but also variations functionally equivalent derived from organisms other than a human (such as a mouse, a hamster, or a dog). Further, it includes polypeptides prepared using polynucleotides obtained by artificially modifying their amino acid sequences encoded thereby by genetic engineering techniques, on the basis of polynucleotides encoding these native polypeptides (i.e., human variations or variations functionally equivalent derived from organisms other than a human), or on the basis of polynucleotides encoding the amino acid sequence of SEQ ID NO: 2 or 4. The term "variation" as used herein means an individual difference between the same polypeptides in the same species or a difference between homologous polypeptides in several species.

[0023]

The variation functionally equivalent derived from a human or organisms other than a human may be obtained by those skilled in the art in accordance with the information of the base sequence of SEQ ID NO: 1 or 3 (for example, the sequence consisting of the 14th to 1882nd bases in the base sequence of SEQ ID NO: 1, or the sequence consisting of the 14th to 1825th bases in the base sequence of SEQ ID NO: 3). For example, an appropriate probe or appropriate primers are designed in accordance with the information of the base sequence. A polymerase chain reaction (PCR) method (Saiki, R. K. et al., Science, 239, 487-491, 1988) or a hybridization method is carried out using a sample (for example, total RNA or an mRNA fraction, a cDNA library, or a phage library) prepared from an organism (for example, a mammal such as a human, a mouse, a hamster, or a dog) of interest and the primers or the probe to obtain a polynucleotide encoding the polypeptide. A desired polypeptide may be obtained by expressing the resulting

polynucleotide in an appropriate expression system, and then, for example, by confirming that the expressed polypeptide exhibits the sodium-calcium exchange activity by a method described in Example 4, and further confirming that the activity is dependent on a potassium ion by a method described in Example 4. In this connection, genetic engineering techniques may be performed in accordance with known methods (for example, Maniatis, T. et al., "Molecular Cloning-A Laboratory Manual", Cold Spring Harbor Laboratory, NY, 1982), unless otherwise specified.

[0024]

Further, the above artificially-modified polypeptide using a polynucleotide modified artificially by genetic engineering techniques may be obtained by, for example, the following procedure. A polynucleotide encoding the polypeptide is obtained by a conventional method such as site-specific mutagenesis (Mark, D. F. et al., Proc. Natl. Acad. Sci. USA, 81, 5662-5666, 1984). A desired polypeptide may be obtained by expressing the resulting polynucleotide in an appropriate expression system, and then, for example, by confirming that the expressed polypeptide exhibits the sodium-calcium exchange activity by a method described in Example 4, and further confirming that the activity is dependent on a potassium ion by a method described in Example 4.

[0025]

The polynucleotide of the present invention is not particularly limited, so long as it encodes the polypeptide of the present invention. As the polynucleotide of the present invention, there may be mentioned, for example, a polynucleotide having the sequence consisting of the 14th to 1882nd bases in the base sequence of SEQ ID NO: 1, or the sequence consisting of the 14th to 1825th bases in the base sequence of SEQ ID NO: 3. In this connection, the term "polynucleotide" as used herein includes both DNA and RNA.

[0026]

A method for producing the polynucleotide of the present invention is not particularly limited, but there may be mentioned, for example, (1) a method using PCR, (2) a

method using conventional genetic engineering techniques (i.e., a method for selecting a transformant comprising a desired cDNA from strains transformed with a cDNA library), or (3) a chemical synthesis method, as described in WO02/052000. These methods will be explained in this order hereinafter.

[0027]

In the method using PCR, the polynucleotide of the present invention may be produced, for example, by the following procedure.

mRNA is extracted from human cells or tissue capable of producing the polypeptide of the present invention. A pair of primers, between which full-length mRNA corresponding to the polypeptide of the present invention or a partial region of the mRNA is located, is synthesized on the basis of the base sequence of a polynucleotide encoding the polypeptide of the present invention. Full-length cDNA encoding the polypeptide of the present invention or a part of the cDNA may be obtained by performing a reverse transcriptase-polymerase chain reaction (RT-PCR) using the extracted mRNA as a template. Further, a desired DNA fragment may be obtained by digesting the obtained DNA with a restriction enzyme or the like and performing ligation, if desired.

[0028]

In the method using conventional genetic engineering techniques, the polynucleotide of the present invention may be produced, for example, by the following procedure.

First, single-stranded cDNA is synthesized by using reverse transcriptase from mRNA prepared by the above-mentioned PCR method as a template, and then double-stranded cDNA is synthesized from the single-stranded cDNA.

Next, a recombinant plasmid comprising the double-stranded cDNA is prepared and introduced into an *Escherichia coli* strain, such as DH 5 α , thereby transforming the strain. A transformant is selected using a drug resistance against, for example, tetracycline or ampicillin as a marker. As a method for selecting a transformant containing the cDNA of interest from the resulting transformants, various methods, such as a method for screening a transformant using a

synthetic oligonucleotide probe or a method for screening a transformant using a probe produced by PCR, may be used.

A method for collecting the polynucleotide of the present invention from the resulting transformant of interest can be carried out in accordance with a known method (for example, Maniatis, T. et al., "Molecular Cloning-A Laboratory Manual", Cold Spring Harbor Laboratory, NY, 1982). For example, it may be carried out by separating a fraction corresponding to the plasmid DNA from cells and cutting out the cDNA region from the plasmid DNA.

[0029]

In the chemical synthesis method, the polynucleotide of the present invention may be produced, for example, by binding DNA fragments produced by a chemical synthesis method. Each DNA fragment can be synthesized using a DNA synthesizer [for example, Oligo 1000M DNA Synthesizer (Beckman) or 394 DNA/RNA Synthesizer (Applied Biosystems)].

In this connection, codons for each amino acid are known and can be optionally selected and determined by the conventional method, for example, by taking a codon usage of each host to be used into consideration (Crantham, R. et al., Nucleic Acids Res., 9, r43-r74, 1981). Further, a partial modification of codons of these base sequences can be carried out in accordance with a conventional method, such as site specific mutagenesis using a primer comprised of a synthetic oligonucleotide encoding for a desired modification (Mark, D. F. et al., Proc. Natl. Acad. Sci. USA, 81, 5662-5666, 1984).

[0030]

Determination of the DNA sequences obtained by the above-mentioned methods can be carried out by, for example, a dideoxynucleotide chain termination method (Messing, J. and Vieira, J., Gene, 19, 269-276, 1982). For example, fluorescence labeled dideoxynucleotides are incorporated to a DNA fragment by a PCR method using a reaction solution containing fluorescence labeled dideoxynucleotides. The amplified DNA fragment is electrophoresed in a sequencer (for example, 3700DNA sequencer; PE Biosystems) and the base sequence thereof can be determined by detecting the

fluorescence.

[0031]

The isolated polynucleotide of the present invention is re-integrated into an appropriate vector DNA and a host cell (including a eucaryotic host cell and a procaryotic host cell) may be transformed by the resulting expression vector. Further, it is possible to express the polynucleotide in a desired host cell, by introducing an appropriate promoter and a sequence related to the gene expression into the vector.

[0032]

The expression vector of the present invention is not particularly limited, so long as it comprises the polynucleotide of the present invention. As the expression vector, there may be mentioned, for example, an expression vector obtained by introducing the polynucleotide of the present invention into a known expression vector appropriately selected in accordance with a host cell to be used.

[0033]

The cell of the present invention is not particularly limited, so long as it is transfected with the expression vector of the present invention and comprises the polynucleotide of the present invention. The cell of the present invention may be, for example, a cell in which the polynucleotide is integrated into a chromosome of a host cell, or a cell containing the polynucleotide as an expression vector comprising the polynucleotide. Further, the cell of the present invention may be a cell expressing the polypeptide of the present invention, or a cell not expressing the polypeptide of the present invention. The cell of the present invention may be obtained by, for example, transfecting a desired host cell with the expression vector of the present invention.

[0034]

In the eucaryotic host cells, for example, cells of vertebrates, insects, and yeast are included. As the vertebral cell, there may be mentioned, for example, a simian COS cell (Gluzman, Y., Cell, 23, 175-182, 1981), a

dihydrofolate reductase defective strain of a Chinese hamster ovary cell (CHO-dhfr⁻ cell) (Urlaub, G. and Chasin, L. A., Proc. Natl. Acad. Sci. USA, 77, 4216-4220, 1980), a Chinese hamster lung fibroblast (Dede cell, ATCC: CCL-39) used in Example 4, a human fetal kidney derived HEK293 cell (ATCC: CRL-1573) used in Example 4, a 293-EBNA cell (Invitrogen) obtained by introducing an EBNA-1 gene of Epstein Barr Virus into HEK293 cell, or L929 cell (ATCC: CRL-2148).

[0035]

As an expression vector for a vertebral cell, a vector containing a promoter positioned upstream of the gene to be expressed, an RNA splicing site, a polyadenylation site, a transcription termination sequence, and the like may be generally used. The vector may further contain a replication origin, if necessary. As the expression vector, there may be mentioned, for example, pSV2dhfr containing an SV40 early promoter (Subramani, S. et al., Mol. Cell. Biol., 1, 854-864, 1981), pEF-BOS containing a human elongation factor promoter (Mizushima, S. and Nagata, S., Nucleic Acids Res., 18, 5322, 1990), pCEP4 containing a cytomegalovirus promoter (Invitrogen), pIRESneo2 (CLONTECH), or pcDNA3.1 (Invitrogen).

[0036]

When the COS cell is used as the host cell, a vector which has an SV40 replication origin, can perform an autonomous replication in the COS cell, and has a transcription promoter, a transcription termination signal, and an RNA splicing site, may be used as the expression vector. As the vector, there may be mentioned, for example, pME18S (Maruyama, K. and Takebe, Y., Med. Immunol., 20, 27-32, 1990), pEF-BOS (Mizushima, S. and Nagata, S., Nucleic Acids Res., 18, 5322, 1990), or pCDM8 (Seed, B., Nature, 329, 840-842, 1987).

[0037]

The expression vector may be incorporated into COS cells by, for example, a DEAE-dextran method (Luthman, H. and Magnusson, G., Nucleic Acids Res., 11, 1295-1308, 1983), a calcium phosphate-DNA co-precipitation method (Graham, F.

L. and van der Ed, A. J., Virology, 52, 456-457, 1973), a method using a commercially available transfection reagent (for example, FuGENE™6 Transfection Reagent; Roche Diagnostics), or an electroporation method (Neumann, E. et al., EMBO J., 1, 841-845, 1982).

[0038]

When the CHO cell is used as the host cell, a transformant capable of stably producing the polypeptide of the present invention can be obtained by carrying out a co-transfection of an expression vector comprising the polynucleotide of the present invention, together with a vector capable of expressing a neo gene which functions as a G418 resistance marker, such as pRSVneo (Sambrook, J. et al., "Molecular Cloning-A Laboratory Manual", Cold Spring Harbor Laboratory, NY, 1989) or pSV2-neo (Southern, P. J. and Berg, P., J. Mol. Appl. Genet., 1, 327-341, 1982), and selecting a G418 resistant colony.

[0039]

When the 293-EBNA cell is used as the host cell, for example, pCEP4 (Invitrogen) containing a replication origin of Epstein Barr Virus and capable of performing an autonomous replication in the 293-EBNA cell may be used as the expression vector.

[0040]

The cell of the present invention may be cultured in accordance with the conventional method, and the polypeptide of the present invention is produced at the surface of the cell. As a medium to be used in the culturing, a medium commonly used in a desired host cell may be appropriately selected. In the case of the COS cell or Dede cell, for example, a medium such as an RPMI-1640 medium or a Dulbecco's modified Eagle's minimum essential medium (DMEM) may be used, by supplementing it with a serum component such as fetal bovine serum (FBS) if necessary. In the case of the 293-EBNA cell, a medium such as a Dulbecco's modified Eagle's minimum essential medium (DMEM) with a serum component such as fetal bovine serum (FBS) and G418 may be used.

[0041]

The polypeptide of the present invention produced at the cell surface by culturing the cell of the present invention may be separated and purified therefrom by various known separation techniques making use of the physical properties, chemical properties and the like of the polypeptide. More particularly, a cell membrane fraction containing the polypeptide of the present invention may be obtained by culturing cells expressing the polypeptide of the present invention, suspending them in a buffer, homogenizing them, and centrifuging the resulting homogenate. After the obtained cell membrane fraction was solubilized, the polypeptide of the present invention may be purified by a commonly used treatment, for example, a treatment with a protein precipitant, ultrafiltration, various liquid chromatography techniques such as molecular sieve chromatography (gel filtration), adsorption chromatography, ion exchange chromatography, affinity chromatography, or high performance liquid chromatography (HPLC), or dialysis, or a combination thereof.

[0042]

2. Screening of therapeutic agents for cell injury due to postischemic reperfusion and an inflammatory disease

A substance which inhibits the polypeptide of the present invention can be screened by using the cell expressing the polypeptide of the present invention so as to exhibit the potassium-dependent sodium-calcium exchange activity.

As described in BACKGROUND ART, it is known that the intracellular calcium overload due to reperfusion plays an important role in the generation of a reperfusion injury and that the potassium-dependent sodium-calcium exchanger is involved in the calcium influx into the cell accompanied by the extrusion of intracellular sodium (reverse direction). Further, it is known that the sodium-calcium exchanger extrudes intracellular sodium outside the cell and incorporates calcium into the cell during postischemic reperfusion, to return the intracellular sodium overload caused by reperfusion to the original state. Furthermore, it is known that an expression of adhesive factors in

leukocytes was increased by an increased intracellular calcium concentration, and that monoclonal antibodies against the adhesive factors suppressed the leukocyte accumulation and reduced the area of infarction. Therefore, it is considered that an inhibition of an intracellular calcium overload in leukocytes suppresses the leukocyte accumulation and leukocyte activation by adhesive factors due to the overload, and has an activity of avoiding cell injury caused by the phenomenon of no reflow.

The polypeptide of the present invention consisting of the sequence of SEQ ID NO: 2 or 4 is a potassium-dependent sodium-calcium exchanger expressed abundantly in peripheral leukocytes. Therefore, it is considered that an agent inhibiting or suppressing the potassium-dependent sodium-calcium exchange (reverse direction) in the polypeptide of the present invention suppresses calcium influx into the cell and inhibits the leukocyte activation or adhesion, and thus is useful in the treatment of a cell injury due to postischemic reperfusion.

[0043]

Further, with respect to the leukocyte activation, postischemic reperfusion and inflammation have common aspects, such as an increased intracellular pH in leukocytes and activation of the sodium-hydrogen exchanger thereby, and an increased intracellular calcium concentration thereby, and thus it is considered that the sodium-calcium exchanger in leukocytes is involved in a leukocyte activation during postischemic reperfusion injury and inflammation. That is, it is considered that an inhibition of sodium-calcium exchanger activation in leukocytes suppresses the increased intracellular calcium concentration in leukocytes and inhibits the leukocyte activation.

[0044]

Therefore, the cell of the present invention per se may be used as a screening tool for an inhibitor of the polypeptide of the present invention and an inhibitor of leukocyte activation (particularly a therapeutic agent for cell injury due to postischemic reperfusion and/or an inflammatory disease).

The term "inhibiting or suppressing the polypeptide of the present invention" means inhibiting or suppressing the potassium-dependent sodium-calcium exchange activity, including the case of inhibiting or suppressing the potassium-dependent sodium-calcium exchange activity by inhibiting or suppressing the expression of the polypeptide.

[0045]

The method for screening an inhibitor of the polypeptide of the present invention and an inhibitor of leukocyte activation (particularly a therapeutic agent for cell injury due to postischemic reperfusion and/or an inflammatory disease) comprises the steps of: bringing the cell of the present invention into contact with a substance to be tested, analyzing whether or not the potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide.

[0046]

Substances to be tested which may be applied to the screening method of the present invention are not particularly limited, but there may be mentioned, for example, various known compounds (including peptides) registered in chemical files, compounds obtained by combinatorial chemistry techniques (Terrett, N. K. et al., Tetrahedron, 51, 8135-8137, 1995), or random peptides prepared by employing a phage display method (Felici, F. et al., J. Mol. Biol., 222, 301-310, 1991) or the like. In addition, culture supernatants of microorganisms, natural components derived from plants or marine organisms, or animal tissue extracts may be used as the test substances for screening. Further, compounds (including peptides) obtained by chemically or biologically modifying compounds (including peptides) selected by the screening method of the present invention may be used.

[0047]

The screening method of the present invention is not particularly limited, so long as it comprises the steps of:

bringing a cell expressing the polypeptide of the present invention so as to function as the potassium-dependent sodium-calcium exchanger (i.e., including a cell which was transfected with an expression vector comprising a polynucleotide encoding the polypeptide of the present invention and expresses the polypeptide so as to function as the potassium-dependent sodium-calcium exchanger, and a naturally occurring cell which expresses the polypeptide of the present invention so as to function as the potassium-dependent sodium-calcium exchanger) into contact with a substance to be tested,

(2) analyzing whether or not the potassium-dependent sodium-calcium exchange activity in the polypeptide is inhibited, and

(3) selecting a substance which inhibits the potassium-dependent sodium-calcium exchange activity in the polypeptide. There may be mentioned, on the basis of differences in methods used for analyzing the inhibition of the polypeptide of the present invention, for example,

[1] a screening method utilizing a radioisotope $^{45}\text{Ca}^{2+}$ ion uptake, or

[2] a screening method utilizing a calcium sensitive dye.

Among these methods, the screening method utilizing a radioisotope $^{45}\text{Ca}^{2+}$ ion uptake is preferable. As the cell used in the screening, the cell of the present invention prepared by transfecting it with an expression vector comprising a polynucleotide encoding the polypeptide of the present invention is preferable.

[0048]

According to the above method [1], a substance which inhibits the polypeptide of the present invention and is useful in treating cell injury due to postischemic reperfusion and/or an inflammatory disease may be screened by the following method. Sodium is incorporated into cells expressing the polypeptide of the present invention at the cell surface, using an ionophore (such as monensin) to a univalent cation; the extracellular solution is changed to a radioisotope $^{45}\text{Ca}^{2+}$ ion-containing extracellular solution with or without a test compound; and an intracellular $^{45}\text{Ca}^{2+}$

radioactivity is measured. Then, whether or not the polypeptide of the present invention is inhibited is analyzed, on the basis of the difference in amounts of radioactivity incorporated into cells in the presence or absence of the test substance. That is, the screening method [1] of the present invention comprises the steps of: incorporating radioisotope $^{45}\text{Ca}^{2+}$ ions into the cell expressing the polypeptide of the present invention at the cell surface, and simultaneously, bringing the cell into contact with a substance to be tested, and detecting an amount of radioactivity incorporated into the cell.

[0049]

For example, the cells expressing the polypeptide of the present invention are treated with an extracellular solution containing a test compound and monensin, to incorporate sodium into the cells. The extracellular solution is changed to an extracellular solution containing the test compound and $^{45}\text{Ca}^{2+}$, to incorporate $^{45}\text{Ca}^{2+}$ into the cells by the sodium-calcium exchange activity. The cells are washed with a solution containing lanthanum, an inhibitor of the sodium-calcium exchange activity, to remove $^{45}\text{Ca}^{2+}$ not incorporated. When the reverse sodium-calcium exchange activity is inhibited, an amount of $^{45}\text{Ca}^{2+}$ influx into the cells is decreased, and thus whether or not the polypeptide of the present invention is inhibited can be analyzed on the basis of the radioactivity in the cells as an indicator of the reverse sodium-calcium exchange activity. More particularly, it is preferable to detect the reverse sodium-calcium exchange activity by the method described in Example 4. An inhibitor of the polypeptide of the present invention can be screened by analyzing a change of radioactivity incorporated into the cells by adding the test substance.

[0050]

When screening a substance which inhibits the polypeptide of the present invention and is useful in treating a cell injury due to postischemic reperfusion and/or an inflammatory disease, by the above method [2], a

calcium sensitive dye is incorporated into the cells expressing the polypeptide of the present invention at the cell surface, and then whether or not the polypeptide of the present invention is inhibited is analyzed, on the basis of a change of a fluorescence intensity thereof in the cells in the presence or absence of the test compound. That is, in the screening method [2] of the present invention, the cells expressing the polypeptide of the present invention at the cell surface are treated with an extracellular solution containing an ionophore (such as monensin) to a univalent cation, to incorporate sodium into the cells. The extracellular solution is changed to an extracellular solution containing a test compound and a calcium sensitive dye, to incorporate the calcium sensitive dye into the cells by the sodium-calcium exchange activity. The cells are washed with a solution containing an inhibitor of the sodium-calcium exchange activity such as lanthanum, to remove the calcium sensitive dye not incorporated. When the reverse sodium-calcium exchange activity is inhibited, a fluorescence intensity of the calcium sensitive dye in the cells is decreased, and thus whether or not the polypeptide of the present invention is inhibited can be analyzed, on the basis of the fluorescence intensity in the cells as an indicator of the reverse sodium-calcium exchange activity. An inhibitor of the polypeptide of the present invention can be screened by detecting the fluorescence intensity of the calcium sensitive dye in the presence of the test compound. The screening method comprises the steps of: incorporating a calcium sensitive dye into the cell, and then bringing the cell into contact with a substance to be tested, and detecting the fluorescence intensity of the calcium sensitive dye in the cell. This screening utilizes the feature that a calcium sensitive dye can optically detect calcium influx accompanied by the reverse-mode activation of sodium-calcium exchanger.

[0051]

More particularly, the activity of the polypeptide of the present invention can be detected using, for example,

Fura-2 or a derivative thereof as the calcium sensitive dye. An inhibitor of the polypeptide of the present invention can be screened by comparing a change of the fluorescence intensity of the dye in the presence or absence of the test substance. When the reverse sodium-calcium exchange activity is inhibited, the fluorescence intensity is decreased.

[0052]

As described above, according to the screening method of the present invention, a substance (particularly a compound) which inhibits the potassium-dependent sodium-calcium exchange activity of the polypeptide of the present invention, i.e., a substance (particularly a compound) having an inhibitory activity, can be screened. A compound having an inhibitory activity to be selected in the screening method of the present invention may be defined as a compound which inhibits an activity of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4. As the compound, a compound having IC₅₀ of 100 $\mu\text{mol/L}$ or less is preferable. For example, when a test compound is reacted for a predetermined period under the conditions described in Example 6, a compound having IC₅₀ of 100 $\mu\text{mol/L}$ or less can be selected as a substance having an inhibitory activity. Using an inhibitory compound isolated by the screening as a main component, a medicament which targets the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4 can be obtained. For example, IC₅₀ of 2-[2-[4-(4-nitrobenzyloxy)phenyl]ethyl]isothiourea methanesulfonate (hereinafter referred to as compound A) selected in Example 6 were 15.8 $\mu\text{mol/L}$ and 35.0 $\mu\text{mol/L}$, with respect to the novel potassium-dependent sodium-calcium exchangers consisting of the amino acid sequences of SEQ ID NOS: 2 and 4, respectively. Further, IC₅₀ of 3',4'-dichlorobenzamil (hereinafter referred to as compound B) were 20.2 $\mu\text{mol/L}$ and 58.9 $\mu\text{mol/L}$, with respect to the novel potassium-dependent sodium-calcium exchangers consisting of the amino acid sequences of SEQ ID NOS: 2 and 4, respectively. From the results, it is apparent that an

inhibitor of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4 can be selected by the screening method of the present invention.

[0053]

3. Medicament of the present invention

The medicament of the present invention comprising, as an active ingredient, a compound (preferably the above compound A or B) which modifies an activity of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4 may be prepared using carriers, fillers, or other additives generally used in the preparation of medicaments, in accordance with the active ingredient.

[0054]

Examples of administration include oral administration by tablets, pills, capsules, granules, fine granules, powders, oral solutions and the like, and parenteral administration by injections (e.g., intravenous, intramuscular, or the like), suppositories, transdermal preparations, transmucosal absorption preparations and the like. Particularly, in the case of peptides which are digested in the stomach, a parenteral administration such as an intravenous injection or the like is preferable.

In the solid composition of the present invention for use in the oral administration, one or more active substances may be mixed with at least one inert diluent such as lactose, mannitol, glucose, microcrystalline cellulose, hydroxypropylcellulose, starch, polyvinyl pyrrolidone, or aluminum magnesium silicate. In the usual way, the composition may contain additives other than the inert diluent, such as a lubricant, a disintegrating agent, a stabilizing agent, or a solubilizing or solubilization assisting agent. If necessary, tablets or pills may be coated with a sugar coating or a film of a gastric or enteric substance.

[0055]

The liquid composition for oral administration may include, for example, emulsions, solutions, suspensions,

syrups, and elixirs, and may contain a generally used inert diluent such as purified water or ethyl alcohol. The composition may contain additives other than the inert diluent, such as moistening agents, suspending agents, sweeteners, flavors, or antiseptics.

The injections for parenteral administration may include aseptic aqueous or non-aqueous solutions, suspensions, and emulsions. Examples of the diluent for use in the aqueous solutions and suspensions include distilled water for injection use and physiological saline. Examples of the diluent for use in the non-aqueous solutions and suspensions include alcohols (e.g., ethanol), glycols (e.g., propylene glycol or polyethylene glycol), polysorbate 80 (trade name) and the like. Such a composition may further contain a moistening agent, an emulsifying agent, a dispersing agent, a stabilizing agent, a solubilizing or solubilization assisting agent, an antiseptic or the like. These compositions may be sterilized, for example, by filtration through a bacteria retaining filter, blending of a germicide, or irradiation. Alternatively, they may be used by first making them into sterile solid compositions and dissolving them in sterile water or other sterile solvent for injection use prior to their use.

[0056]

The dose is optionally decided by taking into consideration the strength of each active ingredient selected by the aforementioned screening method, or symptoms, age, sex, or the like of each patient to be administered. For example, in the case of oral administration, the usual dosage for an adult (60 kg in weight) is about 0.1 to 100 mg, preferably 0.1 to 50 mg per day. In the case of parenteral administration, the usual dosage is about 0.01 to 50 mg, preferably 0.01 to 10 mg per day in the form of an injection.

[0057]

[EXAMPLES]

The present invention now will be further illustrated by, but is by no means limited to, the following Examples. The procedures were performed in accordance with the known methods (for example, Maniatis, T., et al., "Molecular

Cloning - A Laboratory Manual", Cold Spring Harbor Laboratory, NY, 1982; and Hille, B., Ionic Channels of Excitable Membranes, 2nd Ed., Sinauer Associates Inc., MA, 1992), unless otherwise specified.

[0058]

Example 1: Isolation of genes encoding novel potassium-dependent sodium-calcium exchangers and construction of expression vectors

A full-length cDNA encoding each novel sodium-calcium exchanger of the present invention having the amino acid sequence of SEQ ID NO: 2 or 4 was obtained by a reverse transcriptase-polymerase chain reaction (RT-PCR) method using human brain cDNA (Marathon-Ready cDNA; Clontech) as a template, by the following procedure.

[0059]

A PCR was carried out using human brain cDNA (Marathon-Ready cDNA; Clontech) as a template, an oligonucleotide consisting of the base sequence of SEQ ID NO: 5 (having the EcoRI recognition sequence added to the 5'-terminus) as a forward primer, an oligonucleotide consisting of the base sequence of SEQ ID NO: 6 (having the KpnI recognition sequence added to the 5'-terminus) as a reverse primer, and DNA Polymerase (PLATINUM Taq DNA Polymerase High-Fidelity; GIBCO-BRL). In the PCR, a thermal denaturation was first performed at 95°C for 1 minute, and then a cycle consisting of reactions at 98°C for 10 seconds, at 60°C for 20 seconds, and at 72°C for 3 minutes was repeated 40 times. As a result, two DNA bands of approximately 1.9 kbp were amplified. The longer DNA fragment was designated "622" and the shorter one was designated "603".

[0060]

Each DNA fragment was digested with restriction enzymes EcoRI and KpnI, and cloned into plasmid pcDNA3.1 (Invitrogen). The resulting clones were designated pcDNA-622 and pcDNA-603. In this connection, the plasmid pcDNA3.1 contains a cytomegalovirus promoter sequence and may be used for expressing a novel potassium-dependent sodium-calcium exchanger in an animal cell.

[0061]

The base sequences of the clones pcDNA-622 and pcDNA-603 were analyzed using a DNA sequencer (ABI377 DNA Sequencer; Applied Biosystems) by a dideoxy terminator method to obtain the base sequences of SEQ ID NOS: 1 and 3, respectively.

The base sequence of SEQ ID NO: 1 (total base pairs = 1902 bp) contains an open reading frame represented by the sequence consisting of the 14th to 1882nd bases. The amino acid sequence deduced from the open reading frame and consisting of 622 amino acid residues was that of SEQ ID NO: 2.

The base sequence of SEQ ID NO: 3 (total base pairs = 1845 bp) contains an open reading frame represented by the sequence consisting of the 14th to 1825th bases. The amino acid sequence deduced from the open reading frame and consisting of 603 amino acid residues was that of SEQ ID NO: 4.

[0062]

Example 2: Analysis of expression distribution of potassium-dependent sodium-calcium exchangers in human tissues

An expression distribution of the gene (SEQ ID NO: 1) encoding the novel potassium-dependent sodium-calcium in human tissues was analyzed by a reverse transcriptase-polymerase chain reaction (RT-PCR) method in accordance with the following procedure.

Poly A⁺ RNA (5 ng, respectively; Clontech) from each human tissue was treated with DNase, and then a first-strand cDNA was synthesized by carrying out a reverse transcription using an RT-PCR kit (SUPERScript First-Strand Synthesis System for RT-PCR; GIBCO-BRL).

A PCR was carried out using the resulting first-strand cDNA as a template, an oligonucleotide consisting of the base sequence of SEQ ID NO: 7 as a forward primer, an oligonucleotide consisting of the base sequence of SEQ ID NO: 8 as a reverse primer, and DNA Polymerase (PLATINUM Taq DNA Polymerase High-Fidelity; GIBCO-BRL). In the PCR, a thermal denaturation was first performed at 94°C for 1 minute, and then a cycle consisting of reactions at 98°C for 10 seconds, at 64°C for 20 seconds, and at 68°C for 1 minute

and 30 seconds was repeated 35 times. In this connection, the base sequences of the primers are specific sequences commonly contained in both genes encoding the polypeptides consisting of the amino acid sequences of SEQ ID NOS: 2 and 4.

[0063]

When the RT-PCR analysis of human peripheral leukocytes was carried out, DNA fragments of approximately 750 bp and approximately 700 bp were amplified. The DNA fragments of approximately 750 bp and approximately 700 bp contained "the sequence consisting of the 348th to 1101st bases in the base sequence of SEQ ID NO: 1" and "the sequence consisting of the 348th to 1044th bases in the base sequence of SEQ ID NO: 3", respectively. From the result, it was found that the mRNA of each potassium-dependent sodium-calcium exchanger of the present invention was expressed in human peripheral leukocytes.

[0064]

Example 3: Expression of potassium-dependent sodium-calcium exchangers in animal cells HEK293, CCL-39, and CHO

Each polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4 was expressed in animal cells to detect a novel potassium-dependent sodium-calcium exchange activity of the polypeptide. As the animal cells, an HEK293 cell (ATCC: CRL-1573), a Dede cell (ATCC: CCL-39), and a CHO-dhfr⁻ cell (ATCC: CRL-9096) were used. HEK293 cells, Dede cells, or CHO-dhfr⁻ cells were transfected with the expression vector pcDNA-622 or pcDNA-603 prepared in Example 1 and a commercially available transfection reagent (LipofectAMINE2000; GIBCO-BRL) to express each potassium-dependent sodium-calcium exchanger in each cell. In this connection, the concrete procedure was carried out in accordance with a manual attached to the transfection reagent. Further, a cell transfected with the plasmid pcDNA3.1 was prepared as a control cell in a similar fashion. The resulting transfected cells were used in the following Examples 4 and 5.

[0065]

Example 4: Detection of potassium-dependent sodium-calcium

exchange activity

The sodium-calcium exchange activity was measured using each of the cells prepared in Example 3.

Sodium was incorporated into cells by changing the medium to an extracellular solution containing monensin [i.e., a solution containing 0.01 mmol/L monensin, 1 mmol/L ouabain, 146 mmol/L NaCl, 4 mmol/L KCl, 0.1 mmol/L CaCl₂, 2 mmol/L MgCl₂, 10 mmol/L glucose, 0.1% bovine serum albumin, and 10 mmol/L HEPES-Tris (pH=7.4)] and incubating the cells at 37°C for 30 minutes. The solution was changed to an extracellular solution containing calcium chloride (⁴⁵CaCl₂; 55.5 kBq/mL) [i.e., a solution containing 0.01 mmol/L verapamil, 1 mmol/L ouabain, 150 mmol/L NaCl, 0.1 mmol/L CaCl₂, 2 mmol/L MgCl₂, 10 mmol/L glucose, 0.1% bovine serum albumin, and 10 mmol/L HEPES-Tris (pH=7.4); extracellular solution A], and the cells were allowed to stand at room temperature for 15 minutes to exchange the intracellular sodium ions for the calcium ions.

In addition, an extracellular solution in which choline chloride was substituted for NaCl was used. Further, to examine the dependency on a potassium ion, an extracellular solution containing calcium chloride (⁴⁵CaCl₂; 55.5 kBq/mL) and potassium chloride [i.e., a solution containing 0.01 mmol/L verapamil, 1 mmol/L ouabain, 4 mmol/L KCl, 146 mmol/L choline chloride, 0.1 mmol/L CaCl₂, 2 mmol/L MgCl₂, 10 mmol/L glucose, 0.1% bovine serum albumin, and 10 mmol/L HEPES-Tris (pH=7.4)] was used.

[0066]

The treated cells were washed with a washing solution containing 120 mmol/L choline chloride, 10 mmol/L LaCl₃, and 10 mmol/L HEPES-Tris (pH=7.4) to remove calcium ions not incorporated, and then an intracellular calcium ion radioactivity was measured and analyzed by a liquid scintillation counter. In the HEK293 cells, Dede cells, or CHO-dhfr⁻ cells expressing the polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4, a higher radioactivity was measured when using the extracellular solution in which sodium was replaced with choline. Further, higher radioactivity was measured when using the

extracellular solution containing potassium.

[0067]

As described above, it was confirmed that, in the cells expressing the polypeptide consisting of the amino acid sequence of SEQ ID NO: 2 or 4 of the present invention, intracellular sodium ions can be exchanged for extracellular calcium ions more effectively in the presence of potassium ions, and that the polypeptide of the present invention consisting of the amino acid sequence of SEQ ID NO: 2 or 4 exhibits a potassium-dependent sodium-calcium exchange activity.

[0068]

Example 5: Construction of cell strain stably expressing novel potassium-dependent sodium-calcium exchanger

To construct a stable-expression cell strain, the Dede cells prepared in Example 3 were repetitively subcultured in a DMEM medium containing G-418 (GIBCO-BRL) at a final concentration of 400 mg/L, and the cells were seeded into wells of 96-well plates (5000 cells/well). The potassium-dependent sodium-calcium exchange activity in each well was measured in accordance with the method described in Example 4, to select cells having a potassium-dependent sodium-calcium exchange activity in which a radioactivity in the case of the extracellular solution containing potassium was 5000 cpm/well or more and an S/N ratio was 10 or more. The resulting stable-expression cell strains derived from a Dede cell were used in the following Example 6.

[0069]

Example 6: Screening substances inhibiting novel potassium-dependent sodium-calcium exchange activity utilizing radioisotope calcium (⁴⁵Ca) ion release

In accordance with the method described in Example 4, substances which inhibit an activity of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4 were screened by measuring a calcium ion radioactivity.

[0070]

The stable-expression cells derived from a Dede cell, prepared in Example 5, were seeded into each well of 96-well

plates (5000 cells/well), and an inhibitory activity was determined by measuring a calcium ion radioactivity in accordance with the method described in Example 4, to screen substances which inhibit an activity of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4. After cells were treated with monensin, each solution containing a test compound [i.e., a solution of a test compound dissolved in dimethyl sulfoxide (DMSO)] was added to an extracellular solution [a solution containing 0.01 mmol/L verapamil, 1 mmol/L ouabain, 4 mmol/L KCl, 146 mmol/L choline chloride, 0.1 mmol/L CaCl₂, 2 mmol/L MgCl₂, 10 mmol/L glucose, 0.1% bovine serum albumin, and 10 mmol/L HEPES-Tris (pH=7.4)] containing calcium chloride (⁴⁵CaCl₂; 55.5 kBq/mL), and measured to select substances exhibiting an inhibitory activity of 30% or more.

[0071]

As a result, two compounds which exhibited an activity of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4, i.e., 2-[2-[4-(4-nitrobenzyloxy)phenyl]ethyl] isothiourethane methanesulfonate [compound A; Japanese Unexamined Patent Publication (Kokai) 9-67336 or J Biol Chem. 1996. 271(37): 22391-7.] and 3',4'-dichlorobenzamil [compound B; catalog compound No. B-710, NIMH Chemical Inventory, U.S.A or Proc Natl Acad Sci U S A. 1984 May; 81(10): 3238-42.], were obtained.

[0072]

Example 7: Effect of inhibitors in human peripheral polymorphonuclear leukocytes

The compounds obtained in Example 6 were evaluated using human peripheral polymorphonuclear leukocytes (PMNs). A blood sample was obtained by collecting blood from a healthy adult. A ficoll solution for separation (mono-poly resolving medium; Dainippon Pharmaceutical) was poured into each centrifuge tube, and a fresh blood component was laid on the ficoll solution. Centrifugation was carried out at 400×g at room temperature for 30 minutes to separate the upper layer (monocytes), the intermediate layer (PMNs), and

the precipitate (erythrocytes). Each layer was collected in such a way that they were not mixed with each other, suspended in a phosphate buffered saline (PBS), and recentrifuged. Each collected layer was suspended in an RPMI1640 medium containing 0.2% bovine serum albumin (BSA), and used in the following experiment after counting the number of cells. A portion of the resulting PMN component was stained by a hematoxylin and eosin stain to confirm that neutrophils (approximately 95%) and eosinophils (approximately 5%) were contained.

[0073]

A migration test was carried out using a 24-well disposable chemotaxis chamber Transwell 3 μ m; Corning). Each test solution in which compound A or B (final concentration = 10 μ mol/L, 30 μ mol/L, and 100 μ mol/L) were contained in a medium (0.2% BSA, RPMI1640 medium) containing 100 nmol/L formylmethionyl-leucinyI-phenylalanine (fMLP) was prepared, and 1 mL of each test solution was added to each of the lower wells. The collected PMNs were added to each of the upper wells (2.0×10^6 cells/well; 0.2mL) and incubated at 37°C for 1 hour. The numbers of cells which migrated to the lower wells were counted by a hemacytometer. The test was independently repeated three times.

[0074]

The results are shown in Table 1. Each value shown in Table 1 is a relative value when the value obtained in the case of using DMSO instead of the test compounds is regarded as 100. Further, each value shown in Table 1 is an average and a standard deviation (SD) on the basis of three independent experiments. Data obtained in the case that PMNs were not stimulated by fMLP (BG) are shown in Table 1.

As shown in Table 1, compounds A and B inhibited the migration of PMNs dose-dependently. From the results, it was found that compounds which inhibit an activity of the novel potassium-dependent sodium-calcium exchanger consisting of the amino acid sequence of SEQ ID NO: 2 or 4 have an effect capable of inhibiting the migration of PMNs. That is, it was found that such compounds have an advantageous effect as a therapeutic agent for cell injury

due to postischemic reperfusion and/or an inflammatory disease.

[0075]

Table 1

		Number of migrating cells (relative value)
Control	DMSO	100
compound A	10 μ mol/L	78.18 \pm 25.93
	30 μ mol/L	29.50 \pm 26.16
	100 μ mol/L	16.84 \pm 10.78
compound B	10 μ mol/L	100.68 \pm 7.27
	30 μ mol/L	72.90 \pm 15.85
	100 μ mol/L	44.58 \pm 32.82
BG	fMLP (-)	19.22 \pm 6.20

[0076]

[Effects of the Invention]

The polypeptide of the present invention is a potassium-dependent sodium-calcium exchanger expressed in peripheral leukocytes, and thus promotes leukocyte activation, cell injury due to postischemic reperfusion, and inflammation. Therefore, a substance which inhibits the polypeptide of the present invention is useful in treating cell injury due to postischemic reperfusion and/or an inflammatory disease.

Further, the polypeptide of the present invention and the cell of the present invention expressing the polypeptide at the surface thereof are useful in screening an inhibitor of leukocyte activation and a therapeutic agent for cell injury due to postischemic reperfusion and/or an inflammatory disease. A convenient screening system for obtaining an inhibitor of leukocyte activation and a therapeutic agent for cell injury due to postischemic reperfusion and/or an inflammatory disease can be provided by using the cell of the present invention. Furthermore, the polynucleotide and the expression vector of the present invention is useful in manufacturing a screening tool for an inhibitor of leukocyte activation and a therapeutic agent for cell injury and/or an inflammatory disease.

[0077]

According to the screening tool or screening method of the present invention, a substance useful as an inhibitor of leukocyte activation and a therapeutic agent for cell injury

due to postischemic reperfusion and an inflammatory disease can be screened. The pharmaceutical composition of the present invention for inhibiting leukocyte activation is useful in preventing cell injury due to postischemic reperfusion and in treating and/or preventing an inflammatory disease.

[0078]

[Free Text in Sequence Listing]

Features of "Artificial Sequence" are described in the numeric identifier <223> in the Sequence Listing. More particularly, each of the base sequences of SEQ ID NOS: 5 to 8 is an artificially synthesized primer sequence.

[0079]

[Sequence Listing]

<110> Yamanouchi Pharmaceutical Co., Ltd.

<120> Method of screening therapeutic agents for ischemia-reperfusion injury and inflammatory diseases

<130> YAM033286P

<160> 8

<170> PatentIn version 3.1

<210> 1

<211> 1902

<212> DNA

<213> Homo sapiens

 $\langle 220 \rangle$

<221> CDS

<222> (14).. (1882)

<223>

<220>

<223> Inventor: Nozawa, Katsura; Mochizuki Shinobu

<400> 1

gggaattcga tcc atg gcg ctc cgc ggg acc ctc cgg ccg ctc aaa gtt

Met Ala Leu Arg Gly Thr Leu Arg Pro Leu Lys Val

1

5

10

cgc	agg	agg	cga	gag	atg	ctg	ccg	cag	caa	gtc	ggc	ttc	gtg	tgc	gcg	97
Arg	Arg	Arg	Arg	Glu	Met	Leu	Pro	Gln	Gln	Val	Gly	Phe	Val	Cys	Ala	
		15					20					25				
gtg	ctg	gcc	ctg	gtg	tgc	tgt	gcg	tcc	ggc	ctc	ttc	ggc	agc	ttg	ggg	145
Val	Leu	Ala	Leu	Val	Cys	Cys	Ala	Ser	Gly	Leu	Phe	Gly	Ser	Leu	Gly	
		30				35					40					
cac	aaa	aca	gct	tct	gct	agc	aaa	cgt	gtc	ctg	cca	gac	aca	tgg	aga	193
His	Lys	Thr	Ala	Ser	Ala	Ser	Lys	Arg	Val	Leu	Pro	Asp	Thr	Trp	Arg	
45					50					55					60	
aat	aga	aag	ttg	atg	gcc	cca	gtg	aat	ggg	aca	cag	aca	gcc	aag	aac	241
Asn	Arg	Lys	Leu	Met	Ala	Pro	Val	Asn	Gly	Thr	Gln	Thr	Ala	Lys	Asn	
				65				70						75		
tgc	aca	gat	cct	gcg	att	cac	gag	ttc	ccc	aca	gat	ctg	ttc	tcc	aat	289
Cys	Thr	Asp	Pro	Ala	Ile	His	Glu	Phe	Pro	Thr	Asp	Leu	Phe	Ser	Asn	
			80					85					90			
aag	gag	cga	cag	cac	gga	gcc	gtc	ctg	ctg	cac	atc	ctt	ggt	gct	ctg	337
Lys	Glu	Arg	Gln	His	Gly	Ala	Val	Leu	Leu	His	Ile	Leu	Gly	Ala	Leu	
		95					100					105				
tat	atg	ttc	tat	gcc	ttg	gcc	ata	gtg	tgc	gat	gac	ttc	ttt	gtt	ccg	385
Tyr	Met	Phe	Tyr	Ala	Leu	Ala	Ile	Val	Cys	Asp	Asp	Phe	Phe	Val	Pro	
		110				115					120					
tct	cta	gag	aag	atc	tgt	gag	aga	ctc	cat	ctg	agc	gaa	gat	gtg	gct	433
Ser	Leu	Glu	Lys	Ile	Cys	Glu	Arg	Leu	His	Leu	Ser	Glu	Asp	Val	Ala	
125					130					135					140	
gga	gcc	acc	ttc	atg	gct	gca	gga	agc	tca	acg	cca	gag	ctg	ttt	gcg	481
Gly	Ala	Thr	Phe	Met	Ala	Ala	Gly	Ser	Ser	Thr	Pro	Glu	Leu	Phe	Ala	
				145				150						155		
tct	gtt	att	ggg	gtg	ttc	atc	acc	cac	ggg	gac	gtc	ggg	gtg	ggc	acc	529
Ser	Val	Ile	Gly	Val	Phe	Ile	Thr	His	Gly	Asp	Val	Gly	Val	Gly	Thr	
			160					165					170			
atc	gtg	ggc	tct	gct	gtg	ttc	aac	atc	ctg	tgc	ata	att	gga	gtg	tgc	577
Ile	Val	Gly	Ser	Ala	Val	Phe	Asn	Ile	Leu	Cys	Ile	Ile	Gly	Val	Cys	
		175					180					185				
gga	ctg	ttt	gct	ggc	cag	gtg	gtc	cgt	ctg	acg	tgg	tgg	gcc	gtg	tgc	625

Gly	Leu	Phe	Ala	Gly	Gln	Val	Val	Arg	Leu	Thr	Trp	Trp	Ala	Val	Cys	
190						195					200					
cga	gac	tcc	gtg	tac	tac	acc	atc	tct	gtc	atc	gtg	ctc	atc	gtg	ttc	673
Arg	Asp	Ser	Val	Tyr	Tyr	Thr	Ile	Ser	Val	Ile	Val	Leu	Ile	Val	Phe	
205					210					215					220	
ata	tat	gat	gaa	caa	att	gtg	tgg	tgg	gaa	ggc	ctg	gtg	ctc	atc	atc	721
Ile	Tyr	Asp	Glu	Gln	Ile	Val	Trp	Trp	Glu	Gly	Leu	Val	Leu	Ile	Ile	
				225					230						235	
ttg	tat	gtg	ttt	tat	att	ctg	atc	atg	aag	tac	aat	gtg	aag	atg	caa	769
Leu	Tyr	Val	Phe	Tyr	Ile	Leu	Ile	Met	Lys	Tyr	Asn	Val	Lys	Met	Gln	
			240					245					250			
gcc	ttt	ttc	aca	gtc	aaa	caa	aag	agc	att	gca	aac	ggt	aac	ccg	gtc	817
Ala	Phe	Phe	Thr	Val	Lys	Gln	Lys	Ser	Ile	Ala	Asn	Gly	Asn	Pro	Val	
		255					260					265				
aac	agt	gag	ctg	gag	gct	ggt	aat	gat	ttc	tat	gac	ggt	agc	tat	gat	865
Asn	Ser	Glu	Leu	Glu	Ala	Gly	Asn	Asp	Phe	Tyr	Asp	Gly	Ser	Tyr	Asp	
	270					275					280					
gac	cct	tcc	gtg	cca	ttg	ctg	ggg	caa	gtg	aag	gag	aag	cca	cag	tat	913
Asp	Pro	Ser	Val	Pro	Leu	Leu	Gly	Gln	Val	Lys	Glu	Lys	Pro	Gln	Tyr	
285					290					295					300	
ggc	aag	aac	ccc	gtg	gtg	atg	gtg	gac	gag	att	atg	agc	tcc	agc	cct	961
Gly	Lys	Asn	Pro	Val	Val	Met	Val	Asp	Glu	Ile	Met	Ser	Ser	Ser	Pro	
			305					310						315		
ccc	aag	ttc	acc	ttc	cct	gaa	gca	ggc	tta	cga	atc	atg	atc	acc	aat	1009
Pro	Lys	Phe	Thr	Phe	Pro	Glu	Ala	Gly	Leu	Arg	Ile	Met	Ile	Thr	Asn	
			320					325					330			
aag	ttt	gga	ccc	agg	acc	cga	cta	cgg	atg	gcc	agc	agg	atc	atc	att	1057
Lys	Phe	Gly	Pro	Arg	Thr	Arg	Leu	Arg	Met	Ala	Ser	Arg	Ile	Ile	Ile	
		335					340					345				
aat	gag	cgg	cag	aga	ctg	atc	aac	tcg	gcc	aat	ggt	gtg	agc	agt	aag	1105
Asn	Glu	Arg	Gln	Arg	Leu	Ile	Asn	Ser	Ala	Asn	Gly	Val	Ser	Ser	Lys	
		350				355					360					
ccg	ctt	caa	aac	ggg	agg	cac	gag	aac	att	gag	aac	ggg	aat	gtt	cct	1153
Pro	Leu	Gln	Asn	Gly	Arg	His	Glu	Asn	Ile	Glu	Asn	Gly	Asn	Val	Pro	

365	370	375	380	
gtg gaa aac ccc gaa gac cct cag cag aat cag gag cag cag ccg ccg				1201
Val Glu Asn Pro Glu Asp Pro Gln Gln Asn Gln Glu Gln Gln Pro Pro				
	385	390	395	
cca cag cca cca ccg cca gag cca gag ccg gtg gag gct gac ttc ctg				1249
Pro Gln Pro Pro Pro Pro Glu Pro Glu Pro Val Glu Ala Asp Phe Leu				
	400	405	410	
tcc ccc ttc tcc gtg ccg gag gcc aga ggg gac aag gtc aag tgg gtg				1297
Ser Pro Phe Ser Val Pro Glu Ala Arg Gly Asp Lys Val Lys Trp Val				
	415	420	425	
ttc acc tgg ccc ctc atc ttc ctc ctg tgc gtc acc att ccc aac tgc				1345
Phe Thr Trp Pro Leu Ile Phe Leu Leu Cys Val Thr Ile Pro Asn Cys				
	430	435	440	
agc aag ccc ccg tgg gag aag ttc ttc atg gtc acc ttc atc acc gcc				1393
Ser Lys Pro Arg Trp Glu Lys Phe Phe Met Val Thr Phe Ile Thr Ala				
	445	450	455	460
acg ctg tgg atc gct gtg ttc tcc tac atc atg gtg tgg ctg gtg act				1441
Thr Leu Trp Ile Ala Val Phe Ser Tyr Ile Met Val Trp Leu Val Thr				
	465	470	475	
att atc gga tac aca ctt ggg atc ccg gat gtc atc atg ggc att act				1489
Ile Ile Gly Tyr Thr Leu Gly Ile Pro Asp Val Ile Met Gly Ile Thr				
	480	485	490	
ttc ctg gca gca ggg aca agt gtt cca gac tgc atg gcc agc cta att				1537
Phe Leu Ala Ala Gly Thr Ser Val Pro Asp Cys Met Ala Ser Leu Ile				
	495	500	505	
gtg gcg aga caa ggc ctt ggg gac atg gca gtc tcc aac acc ata gga				1585
Val Ala Arg Gln Gly Leu Gly Asp Met Ala Val Ser Asn Thr Ile Gly				
	510	515	520	
agc aac gtg ttt gac atc ctg gta gga ctt ggt gta ccg tgg ggc ctg				1633
Ser Asn Val Phe Asp Ile Leu Val Gly Leu Gly Val Pro Trp Gly Leu				
	525	530	535	540
cag acc atg gtt gtt aat tat gga tca aca gtg aag atc aac agc cgg				1681
Gln Thr Met Val Val Asn Tyr Gly Ser Thr Val Lys Ile Asn Ser Arg				
	545	550	555	

ggg ctg gtc tat tcc gtg gtc ctg ttg ctg ggc tct gtc gct ctc acc 1729
 Gly Leu Val Tyr Ser Val Val Leu Leu Leu Gly Ser Val Ala Leu Thr
 560 565 570

gtc ctc ggc atc cac cta aac aag tgg cga ctg gac cgg aag ctg ggt 1777
Val Leu Gly Ile His Leu Asn Lys Trp Arg Leu Asp Arg Lys Leu Gly
575 580 585

gtc tac gtg ctg gtt ctc tac gcc atc ttc ttg tgc ttc tcc ata atg 1825
Val Tyr Val Leu Val Leu Tyr Ala Ile Phe Leu Cys Phe Ser Ile Met
590 595 600

ata gag ttt aac gtc ttt acc ttc gtc aac ttg ccg atg tgc cgg gaa 1873
Ile Glu Phe Asn Val Phe Thr Phe Val Asn Leu Pro Met Cys Arg Glu
605 610 615 620

gac gat tag cgctgagtcg cggtagctgg 1902
Asp Asp

<210>	2
<211>	622
<212>	PRT
<213>	Homo sapiens

.<400> 2

Met Ala Leu Arg Gly Thr Leu Arg Pro Leu Lys Val Arg Arg Arg Arg
1 5 10 15

Glu Met Leu Pro Gln Gln Val Gly Phe Val Cys Ala Val Leu Ala Leu
20 25 30

Val Cys Cys Ala Ser Gly Leu Phe Gly Ser Leu Gly His Lys Thr Ala
35 40 45

Ser Ala Ser Lys Arg Val Leu Pro Asp Thr Trp Arg Asn Arg Lys Leu
50 55 60

Met Ala Pro Val Asn Gly Thr Gln Thr Ala Lys Asn Cys Thr Asp Pro
 65 70 75 80

Ala Ile His Glu Phe Pro Thr Asp Leu Phe Ser Asn Lys Glu Arg Gln
 85 90 95

His Gly Ala Val Leu Leu His Ile Leu Gly Ala Leu Tyr Met Phe Tyr
 100 105 110

Ala Leu Ala Ile Val Cys Asp Asp Phe Phe Val Pro Ser Leu Glu Lys
 115 120 125

Ile Cys Glu Arg Leu His Leu Ser Glu Asp Val Ala Gly Ala Thr Phe
 130 135 140

Met Ala Ala Gly Ser Ser Thr Pro Glu Leu Phe Ala Ser Val Ile Gly
 145 150 155 160

Val Phe Ile Thr His Gly Asp Val Gly Val Gly Thr Ile Val Gly Ser
 165 170 175

Ala Val Phe Asn Ile Leu Cys Ile Ile Gly Val Cys Gly Leu Phe Ala
 180 185 190

Gly Gln Val Val Arg Leu Thr Trp Trp Ala Val Cys Arg Asp Ser Val
 195 200 205

Tyr Tyr Thr Ile Ser Val Ile Val Leu Ile Val Phe Ile Tyr Asp Glu
 210 215 220

Gln Ile Val Trp Trp Glu Gly Leu Val Leu Ile Ile Leu Tyr Val Phe
 225 230 235 240

Tyr Ile Leu Ile Met Lys Tyr Asn Val Lys Met Gln Ala Phe Phe Thr

245	250	255
Val Lys Gln Lys Ser Ile Ala Asn Gly Asn Pro Val Asn Ser Glu Leu		
260	265	270
Glu Ala Gly Asn Asp Phe Tyr Asp Gly Ser Tyr Asp Asp Pro Ser Val		
275	280	285
Pro Leu Leu Gly Gln Val Lys Glu Lys Pro Gln Tyr Gly Lys Asn Pro		
290	295	300
Val Val Met Val Asp Glu Ile Met Ser Ser Ser Pro Pro Lys Phe Thr		
305	310	315
Phe Pro Glu Ala Gly Leu Arg Ile Met Ile Thr Asn Lys Phe Gly Pro		
325	330	335
Arg Thr Arg Leu Arg Met Ala Ser Arg Ile Ile Ile Asn Glu Arg Gln		
340	345	350
Arg Leu Ile Asn Ser Ala Asn Gly Val Ser Ser Lys Pro Leu Gln Asn		
355	360	365
Gly Arg His Glu Asn Ile Glu Asn Gly Asn Val Pro Val Glu Asn Pro		
370	375	380
Glu Asp Pro Gln Gln Asn Gln Glu Gln Gln Pro Pro Pro Gln Pro Pro		
385	390	395
Pro Pro Glu Pro Glu Pro Val Glu Ala Asp Phe Leu Ser Pro Phe Ser		
405	410	415
Val Pro Glu Ala Arg Gly Asp Lys Val Lys Trp Val Phe Thr Trp Pro		
420	425	430

Leu Ile Phe Leu Leu Cys Val Thr Ile Pro Asn Cys Ser Lys Pro Arg
435 440 445

Trp Glu Lys Phe Phe Met Val Thr Phe Ile Thr Ala Thr Leu Trp Ile
450 455 460

Ala Val Phe Ser Tyr Ile Met Val Trp Leu Val Thr Ile Ile Gly Tyr
465 470 475 480

Thr Leu Gly Ile Pro Asp Val Ile Met Gly Ile Thr Phe Leu Ala Ala
485 490 495

Gly Thr Ser Val Pro Asp Cys Met Ala Ser Leu Ile Val Ala Arg Gln
500 505 510

Gly Leu Gly Asp Met Ala Val Ser Asn Thr Ile Gly Ser Asn Val Phe
515 520 525

Asp Ile Leu Val Gly Leu Gly Val Pro Trp Gly Leu Gln Thr Met Val
530 535 540

Val Asn Tyr Gly Ser Thr Val Lys Ile Asn Ser Arg Gly Leu Val Tyr
545 550 555 560

Ser Val Val Leu Leu Leu Gly Ser Val Ala Leu Thr Val Leu Gly Ile
565 570 575

His Leu Asn Lys Trp Arg Leu Asp Arg Lys Leu Gly Val Tyr Val Leu
580 585 590

Val Leu Tyr Ala Ile Phe Leu Cys Phe Ser Ile Met Ile Glu Phe Asn
595 600 605

Val Phe Thr Phe Val Asn Leu Pro Met Cys Arg Glu Asp Asp
610 615 620

$\langle 210 \rangle$	3
$\langle 211 \rangle$	1845
$\langle 212 \rangle$	DNA
$\langle 213 \rangle$	Homo sapiens

<220>
 <221> CDS
 <222> (14).. (1825)
 <223>

[illegible]

cgc agg agg cga gag atg ctg ccg cag caa gtc ggc ttc gtg tgc gcg 97
 Arg Arg Arg Arg Glu Met Leu Pro Gln Gln Val Gly Phe Val Cys Ala
 15 20 25

gtg ctg gcc ctg gtg tgc tgt gcg tcc ggc ctc ttc ggc agc ttg ggg 145
Val Leu Ala Leu Val Cys Cys Ala Ser Gly Leu Phe Gly Ser Leu Gly
30 35 40

cac aaa aca gct tct gct agc aaa cgt gtc ctg cca gac aca tgg aga 193
His Lys Thr Ala Ser Ala Ser Lys Arg Val Leu Pro Asp Thr Trp Arg
45 50 55 60

aat aga aag ttg atg gcc cca gtg aat ggg aca cag aca gcc aag aac 241
Asn Arg Lys Leu Met Ala Pro Val Asn Gly Thr Gln Thr Ala Lys Asn
65 70 75

tgc aca gat cct gcg att cac gag ttc ccc aca gat ctg ttc tcc aat 289
Cys Thr Asp Pro Ala Ile His Glu Phe Pro Thr Asp Leu Phe Ser Asn
80 85 90

aag gag cga cag cac gga gcc gtc ctg ctg cac atc ctt ggt gct ctg 337
Lys Glu Arg Gln His Gly Ala Val Leu Leu His Ile Leu Gly Ala Leu
95 100 105

tat atg ttc tat gcc ttg gcc ata gtg tgc gat gac ttc ttt gtt ccg	385
Tyr Met Phe Tyr Ala Leu Ala Ile Val Cys Asp Asp Phe Phe Val Pro	
110 115 120	
tct cta gag aag atc tgt gag aga ctc cat ctg agc gaa gat gtg gct	433
Ser Leu Glu Lys Ile Cys Glu Arg Leu His Leu Ser Glu Asp Val Ala	
125 130 135 140	
gga gcc acc ttc atg gct gca gga agc tca acg cca gag ctg ttt gcg	481
Gly Ala Thr Phe Met Ala Ala Gly Ser Ser Thr Pro Glu Leu Phe Ala	
145 150 155	
tct gtt att ggg gtg ttc atc acc cac ggg gac gtc ggg gtg ggc acc	529
Ser Val Ile Gly Val Phe Ile Thr His Gly Asp Val Gly Val Gly Thr	
160 165 170	
atc gtg ggc tct gct gtg ttc aac atc ctg tgc ata att gga gtg tgc	577
Ile Val Gly Ser Ala Val Phe Asn Ile Leu Cys Ile Ile Gly Val Cys	
175 180 185	
gga ctg ttt gct ggc cag gtg gtc cgt ctg acg tgg tgg gcc gtg tgc	625
Gly Leu Phe Ala Gly Gln Val Val Arg Leu Thr Trp Trp Ala Val Cys	
190 195 200	
cga gac tcc gtg tac tac acc atc tct gtc atc gtg ctc atc gtg ttc	673
Arg Asp Ser Val Tyr Tyr Thr Ile Ser Val Ile Val Leu Ile Val Phe	
205 210 215 220	
ata tat gat gaa caa att gtg tgg tgg gaa ggc ctg gtg ctc atc atc	721
Ile Tyr Asp Glu Gln Ile Val Trp Trp Glu Gly Leu Val Leu Ile Ile	
225 230 235	
ttg tat gtg ttt tat att ctg atc atg aag tac aat gtg aag atg caa	769
Leu Tyr Val Phe Tyr Ile Leu Ile Met Lys Tyr Asn Val Lys Met Gln	
240 245 250	
gcc ttt ttc aca gtc aaa caa aag agc att gca aac ggt aac ccg gtc	817
Ala Phe Phe Thr Val Lys Gln Lys Ser Ile Ala Asn Gly Asn Pro Val	
255 260 265	
aac agt gag ctg gag gct gtg aag gag aag cca cag tat ggc aag aac	865
Asn Ser Glu Leu Glu Ala Val Lys Glu Lys Pro Gln Tyr Gly Lys Asn	
270 275 280	
ccc gtg gtg atg gtg gac gag att atg agc tcc agc cct ccc aag ttc	913

tac aca ctt ggg atc ccg gat gtc atc atg ggc att act ttc ctg gca 1441
Tyr Thr Leu Gly Ile Pro Asp Val Ile Met Gly Ile Thr Phe Leu Ala

465	470	475	
gca ggg aca agt gtt cca gac tgc atg gcc agc cta att gtg gcg aga			1489
Ala Gly Thr Ser Val Pro Asp Cys Met Ala Ser Leu Ile Val Ala Arg			
480	485	490	
caa ggc ctt ggg gac atg gca gtc tcc aac acc ata gga agc aac gtg			1537
Gln Gly Leu Gly Asp Met Ala Val Ser Asn Thr Ile Gly Ser Asn Val			
495	500	505	
ttt gac atc ctg gta gga ctt ggt gta ccg tgg ggc ctg cag acc atg			1585
Phe Asp Ile Leu Val Gly Leu Gly Val Pro Trp Gly Leu Gln Thr Met			
510	515	520	
gtt gtt aat tat gga tca aca gtg aag atc aac agc cgg ggg ctg gtc			1633
Val Val Asn Tyr Gly Ser Thr Val Lys Ile Asn Ser Arg Gly Leu Val			
525	530	535	540
tat tcc gtg gtc ctg ttg ctg ggc tct gtc gct ctc acc gtc ctc ggc			1681
Tyr Ser Val Val Leu Leu Leu Gly Ser Val Ala Leu Thr Val Leu Gly			
545	550	555	
atc cac cta aac aag tgg cga ctg gac cgg aag ctg ggt gtc tac gtg			1729
Ile His Leu Asn Lys Trp Arg Leu Asp Arg Lys Leu Gly Val Tyr Val			
560	565	570	
ctg gtt ctc tac gcc atc ttc ttg tgc ttc tcc ata atg ata gag ttt			1777
Leu Val Leu Tyr Ala Ile Phe Leu Cys Phe Ser Ile Met Ile Glu Phe			
575	580	585	
aac gtc ttt acc ttc gtc aac ttg ccg atg tgc cgg gaa gac gat tag			1825
Asn Val Phe Thr Phe Val Asn Leu Pro Met Cys Arg Glu Asp Asp			
590	595	600	
cgctgagtcg cggtacctgg			1845

<210> 4
 <211> 603
 <212> PRT
 <213> Homo sapiens

<400> 4

Met Ala Leu Arg Gly Thr Leu Arg Pro Leu Lys Val Arg Arg Arg Arg

1	5	10	15
Glu Met Leu Pro Gln Gln Val Gly Phe Val Cys Ala Val Leu Ala Leu			
	20	25	30
Val Cys Cys Ala Ser Gly Leu Phe Gly Ser Leu Gly His Lys Thr Ala			
	35	40	45
Ser Ala Ser Lys Arg Val Leu Pro Asp Thr Trp Arg Asn Arg Lys Leu			
	50	55	60
Met Ala Pro Val Asn Gly Thr Gln Thr Ala Lys Asn Cys Thr Asp Pro			
65		70	75
Ala Ile His Glu Phe Pro Thr Asp Leu Phe Ser Asn Lys Glu Arg Gln			
	85	90	95
His Gly Ala Val Leu Leu His Ile Leu Gly Ala Leu Tyr Met Phe Tyr			
	100	105	110
Ala Leu Ala Ile Val Cys Asp Asp Phe Phe Val Pro Ser Leu Glu Lys			
	115	120	125
Ile Cys Glu Arg Leu His Leu Ser Glu Asp Val Ala Gly Ala Thr Phe			
	130	135	140
Met Ala Ala Gly Ser Ser Thr Pro Glu Leu Phe Ala Ser Val Ile Gly			
145		150	155
Val Phe Ile Thr His Gly Asp Val Gly Val Gly Thr Ile Val Gly Ser			
	165	170	175
Ala Val Phe Asn Ile Leu Cys Ile Ile Gly Val Cys Gly Leu Phe Ala			
	180	185	190

Gly Gln Val Val Arg Leu Thr Trp Trp Ala Val Cys Arg Asp Ser Val
195 200 205

Tyr Tyr Thr Ile Ser Val Ile Val Leu Ile Val Phe Ile Tyr Asp Glu
210 215 220

Gln Ile Val Trp Trp Glu Gly Leu Val Leu Ile Ile Leu Tyr Val Phe
225 230 235 240

Tyr Ile Leu Ile Met Lys Tyr Asn Val Lys Met Gln Ala Phe Phe Thr
245 250 255

Val Lys Gln Lys Ser Ile Ala Asn Gly Asn Pro Val Asn Ser Glu Leu
260 265 270

Glu Ala Val Lys Glu Lys Pro Gln Tyr Gly Lys Asn Pro Val Val Met
275 280 285

Val Asp Glu Ile Met Ser Ser Ser Pro Pro Lys Phe Thr Phe Pro Glu
290 295 300

Ala Gly Leu Arg Ile Met Ile Thr Asn Lys Phe Gly Pro Arg Thr Arg
305 310 315 320

Leu Arg Met Ala Ser Arg Ile Ile Ile Asn Glu Arg Gln Arg Leu Ile
325 330 335

Asn Ser Ala Asn Gly Val Ser Ser Lys Pro Leu Gln Asn Gly Arg His
340 345 350

Glu Asn Ile Glu Asn Gly Asn Val Pro Val Glu Asn Pro Glu Asp Pro
355 360 365

Gln Gln Asn Gln Glu Gln Gln Pro Pro Pro Gln Pro Pro Pro Pro Glu
370 375 380

Pro Glu Pro Val Glu Ala Asp Phe Leu Ser Pro Phe Ser Val Pro Glu
385 390 395 400

Ala Arg Gly Asp Lys Val Lys Trp Val Phe Thr Trp Pro Leu Ile Phe
405 410 415

Leu Leu Cys Val Thr Ile Pro Asn Cys Ser Lys Pro Arg Trp Glu Lys
420 425 430

Phe Phe Met Val Thr Phe Ile Thr Ala Thr Leu Trp Ile Ala Val Phe
435 440 445

Ser Tyr Ile Met Val Trp Leu Val Thr Ile Ile Gly Tyr Thr Leu Gly
450 455 460

Ile Pro Asp Val Ile Met Gly Ile Thr Phe Leu Ala Ala Gly Thr Ser
465 470 475 480

Val Pro Asp Cys Met Ala Ser Leu Ile Val Ala Arg Gln Gly Leu Gly
485 490 495

Asp Met Ala Val Ser Asn Thr Ile Gly Ser Asn Val Phe Asp Ile Leu
500 505 510

Val Gly Leu Gly Val Pro Trp Gly Leu Gln Thr Met Val Val Asn Tyr
515 520 525

Gly Ser Thr Val Lys Ile Asn Ser Arg Gly Leu Val Tyr Ser Val Val
530 535 540

Leu Leu Leu Gly Ser Val Ala Leu Thr Val Leu Gly Ile His Leu Asn
 545 550 555 560

Lys Trp Arg Leu Asp Arg Lys Leu Gly Val Tyr Val Leu Val Leu Tyr
 565 570 575

Ala Ile Phe Leu Cys Phe Ser Ile Met Ile Glu Phe Asn Val Phe Thr
 580 585 590

Phe Val Asn Leu Pro Met Cys Arg Glu Asp Asp
 595 600

<210> 5

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of artificial Sequence: an artificially synthesized primer sequence

<400> 5

caggaattcc accatggcgc tccgcgggac cctc

34

<210> 6

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of artificial Sequence: an artificially synthesized primer sequence

<400> 6

ccaggtaccg cgactcagcg ctaatcg

27

<210> 7

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of artificial Sequence: an artificially synthesized primer sequence

<400> 7

atgccttggc catagtgtgc gatg

24

<210> 8

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of artificial Sequence: an artificially synthesized primer sequence

<400> 8

ctgctcacac cattggccga gttg

24

[DOCUMENT NAME] Abstract

[ABSTRACT]

[OBJECT] A convenient screening system for obtaining a substance useful in treating cell injuries due to postischemic reperfusion or an inflammatory disease is provided. A novel polypeptide, a polynucleotide encoding the polypeptide, an expression vector comprising the polynucleotide, a cell transfected by the expression vector, and a method for producing the polypeptide, which are useful in constructing the screening system, are provided.

[MEANS FOR SOLUTION] The polypeptide is a potassium-dependent sodium-calcium exchanger expressed in peripheral leukocytes.

[SELECTED DRAWINGS] None